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RAW SEQUENCE LISTING

DATE: 12/26/2001

PATENT APPLICATION: US/09/875,076

TIME: 16:20:14

Input Set : N:\CRF3\RULE60\09875076.RAW.txt

Output Set: N:\CRF3\12262001\I875076.raw

1 <110> APPLICANT: Chen, Ruoping
2 Dang, Huong T.
3 Liaw, Chen W.
4 Lin, I-Lin
5 <120> TITLE OF INVENTION: Human Orphan G Protein Coupled Receptors
6 <130> FILE REFERENCE: AREN0050
7 <140> CURRENT APPLICATION NUMBER: 09/875,076
8 <141> CURRENT FILING DATE: 2001-06-06
9 <150> PRIOR APPLICATION NUMBER: 09/417,044
10 <151> PRIOR FILING DATE: 1999-10-12
11 <150> PRIOR APPLICATION NUMBER: 60/120,416
12 <151> PRIOR FILING DATE: 1999-02-16
13 <150> PRIOR APPLICATION NUMBER: 60/121,851
14 <151> PRIOR FILING DATE: 1999-02-26
15 <150> PRIOR APPLICATION NUMBER: 60/123,946
16 <151> PRIOR FILING DATE: 1999-03-12
17 <150> PRIOR APPLICATION NUMBER: 60/123,949
18 <151> PRIOR FILING DATE: 1999-03-12
19 <150> PRIOR APPLICATION NUMBER: 60/136,436
20 <151> PRIOR FILING DATE: 1999-05-28
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41 <150> PRIOR APPLICATION NUMBER: 60/157,280
42 <151> PRIOR FILING DATE: 1999-10-01
43 <150> PRIOR APPLICATION NUMBER: 60/157,294
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61   agtccattgc ttagatatag ttttgaaacc atggtctccc ctggtttgag ttccttgacc 180
62   gtgaatagta cagctgtgcc cacaacacca gcagcattta agagcctaaa ctgacctctt 240
63   cagatcaccc ttctgctat aatgatattc attctgtttg tgtctttctt tgggaacttg 300
64   qttgtttgcc tcatggttta ccaaaaagct gccatgaggt ctgcaattaa catctcctt 360
65   gccagcctag cttttgcaga catgttgctt gcagtgtcga acatgccctt tgccctgcta 420
66   actattctta ctaccgatg gatttttggg aaattcttct gtagggtatc tgcattgttt 480
67   ttctggttat ttgtgataga aggagtagcc atcctgctca tcattagcat agataggttc 540
68   cttattatag tcagaggcca ggataagcta aacctatata gagctaaggt tctgattgca 600
69   gtttcttggg caacttcctt ttgtgtagct tttcctttag ccgtaggaaa ccccgacctg 660
70   cagataacct cccgagctcc ccagtggtg ttgggttaca caaccaatcc aggtaccag 720
71   gcttatgtga ttttgatttc tctcatttct ttcttcatac ccttctgggt aatactgtac 780
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73   gaaggatatg gctcagccca ggcacagcaa ctgggtctca tgagtctgca gagacctttc 900
74   cagatgagca ttgacatgg ctttaaaaca cgtgccttca ccactatttt gattctcttt 960
75   gctgtcttca ttgtctgtg ggccttcatt accacttaca gcttgtggc aacattcagt 1020
76   aagcactttt actatcagca caactttttt gagattagca cctggctact gtggctctgc 1080
77   tactccaagt ctgcattgaa tccgctgac tactactgga ggattaagaa attccatgat 1140
78   gcttgctgg acatgatgcc taagtcttc aaqgttttgc cgcagctccc tgggtcacaca 1200
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83 <212> TYPE: PRT
84 <213> ORGANISM: Homo sapiens
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88   Thr Thr Phe Val Val Tyr Glu Asn Thr Tyr Met Asn Ile Thr Leu Pro
89             20             25             30
90   Pro Pro Phe Gln His Pro Asp Leu Ser Pro Leu Leu Arg Tyr Ser Phe
91             35             40             45
92   Glu Thr Met Ala Pro Thr Gly Leu Ser Ser Leu Thr Val Asn Ser Thr
93             50             55             60
94   Ala Val Pro Thr Thr Pro Ala Ala Phe Lys Ser Leu Asn Leu Pro Leu
95             65             70             75             80
96   Gln Ile Thr Leu Ser Ala Ile Met Ile Phe Ile Leu Phe Val Ser Phe
97             85             90             95
98   Leu Gly Asn Leu Val Val Cys Leu Met Val Tyr Gln Lys Ala Ala Met

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102  Leu Leu Ala Val Leu Asn Met Pro Phe Ala Leu Val Thr Ile Leu Thr
103          130          135          140
104  Thr Arg Trp Ile Phe Gly Lys Phe Phe Cys Arg Val Ser Ala Met Phe
105          145          150          155          160
106  Phe Trp Leu Phe Val Ile Glu Gly Val Ala Ile Leu Leu Ile Ile Ser
107          165          170          175
108  Ile Asp Arg Phe Leu Ile Ile Val Gln Arg Gln Asp Lys Leu Asn Pro
109          180          185          190
110  Tyr Arg Ala Lys Val Leu Ile Ala Val Ser Trp Ala Thr Ser Phe Cys
111          195          200          205
112  Val Ala Phe Pro Leu Ala Val Gly Asn Pro Asp Leu Gln Ile Pro Ser
113          210          215          220
114  Arg Ala Pro Gln Cys Val Phe Gly Tyr Thr Thr Asn Pro Gly Tyr Gln
115          225          230          235          240
116  Ala Tyr Val Ile Leu Ile Ser Leu Ile Ser Phe Phe Ile Pro Phe Leu
117          245          250          255
118  Val Ile Leu Tyr Ser Phe Met Gly Ile Leu Asn Thr Leu Arg His Asn
119          260          265          270
120  Ala Leu Arg Ile His Ser Tyr Pro Glu Gly Ile Cys Leu Ser Gln Ala
121          275          280          285
122  Ser Lys Leu Gly Leu Met Ser Leu Gln Arg Pro Phe Gln Met Ser Ile
123          290          295          300
124  Asp Met Gly Phe Lys Thr Arg Ala Phe Thr Thr Ile Leu Ile Leu Phe
125          305          310          315          320
126  Ala Val Phe Ile Val Cys Trp Ala Pro Phe Thr Thr Tyr Ser Leu Val
127          325          330          335
128  Ala Thr Phe Ser Lys His Phe Tyr Tyr Gln His Asn Phe Phe Glu Ile
129          340          345          350
130  Ser Thr Trp Leu Leu Trp Leu Cys Tyr Leu Lys Ser Ala Leu Asn Pro
131          355          360          365
132  Leu Ile Tyr Tyr Trp Arg Ile Lys Lys Phe His Asp Ala Cys Leu Asp
133          370          375          380
134  Met Met Pro Lys Ser Phe Lys Phe Leu Pro Gln Leu Pro Gly His Thr
135          385          390          395          400
136  Lys Arg Arg Ile Arg Pro Ser Ala Val Tyr Val Cys Gly Glu His Arg
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143 <213> ORGANISM: Homo sapiens
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146  accaacgcgc tgcacttggt ggtctacagc ttggtgctgg ctgccgggct cccctcaac 120
147  gcgctagccc tctgggtctt cctgcgcgcg ctgcgcgtgc actcgggtgt ggcgtgttac 180
148  atgtgtaacc tggcggccag cgacctgtct ttcacctct cgtgcgccgt tcgtctctcc 240

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151  gccgcacatg  tgcacccgct  ggcactgcgc  cacctgcggc  gggcccgctg  ggcggggctg  420
152  ctctgcctgq  gcctgtgqgc  gctcactcct  gtgtttgcgc  tgcccgccgc  ccgcgtgcac  480
153  aggccctcgc  gttgcgccta  ccgggacctc  gaggtgcgcc  tatgtctcga  gagcttcagc  540
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156  cccgacgcca  cgcagagcca  gcggcgggcg  aagaccgtgc  gcctctgctt  ggctaacctc  720
157  gtcatcttcc  tgcctgtgct  cgtgccctac  aacagcacgc  tgccggctct  cgggctgctg  780
158  cggagcaagg  tgggtggcgg  cagcgtgcct  gcccgcgatc  gcgtgcgcgg  ggtgctgatg  840
159  gtgatgtgtc  tqctggccgg  ccccaactgc  gtgctggacc  cgtgtgtgta  ctactttagc  900
160  gccgagggct  tccgcaaac  cctgcgcggc  ctgggcactc  cgcaccgggc  caggacctcg  960
161  gccaccaacg  ggcgcggggc  ggcgctcgcg  caatccgaaa  ggtccgcctg  caccaccgac  1020
162  gccaccaggc  cggatgcgcg  cagtcagggg  ctgctccgac  cctccgactc  ccactctctg  1080
163  tcttcttcca  cacagtgtcc  ccaggattcc  gccctctga  1119

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166 <211> LENGTH: 372

167 <212> TYPE: PRT

168 <213> ORGANISM: Homo sapiens

169 <400> SEQUENCE: 4

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173      20          25          30
174  Leu  Ala  Ala  Gly  Leu  Pro  Leu  Asn  Ala  Leu  Ala  Leu  Trp  Val  Phe  Leu
175      35          40          45
176  Arg  Ala  Leu  Arg  Val  His  Ser  Val  Val  Ser  Val  Tyr  Met  Cys  Asn  Leu
177      50          55          60
178  Ala  Ala  Ser  Asp  Leu  Leu  Phe  Thr  Leu  Ser  Leu  Pro  Val  Arg  Leu  Ser
179      65          70          75          80
180  Tyr  Tyr  Ala  Leu  His  His  Trp  Pro  Phe  Pro  Asp  Leu  Leu  Cys  Gln  Thr
181      85          90          95
182  Thr  Gly  Ala  Ile  Phe  Gln  Met  Asn  Met  Tyr  Gly  Ser  Cys  Ile  Phe  Leu
183      100         105         110
184  Met  Leu  Ile  Asn  Val  Asp  Arg  Tyr  Ala  Ala  Ile  Val  His  Pro  Leu  Arg
185      115         120         125
186  Leu  Arg  His  Leu  Arg  Arg  Pro  Arg  Val  Ala  Arg  Leu  Leu  Cys  Leu  Gly
187      130         135         140
188  Val  Trp  Ala  Leu  Ile  Leu  Val  Phe  Ala  Val  Pro  Ala  Ala  Arg  Val  His
189      145         150         155         160
190  Arg  Pro  Ser  Arg  Cys  Arg  Tyr  Arg  Asp  Leu  Glu  Val  Arg  Leu  Cys  Phe
191      165         170         175
192  Glu  Ser  Phe  Ser  Asp  Glu  Leu  Trp  Lys  Gly  Arg  Leu  Leu  Pro  Leu  Val
193      180         185         190
194  Leu  Leu  Ala  Glu  Ala  Leu  Gly  Phe  Leu  Leu  Pro  Leu  Ala  Ala  Val  Val
195      195         200         205
196  Tyr  Ser  Ser  Gly  Arg  Val  Phe  Trp  Thr  Leu  Ala  Arg  Pro  Asp  Ala  Thr
197      210         215         220
198  Gln  Ser  Gln  Arg  Arg  Arg  Lys  Thr  Val  Arg  Leu  Leu  Leu  Ala  Asn  Leu

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200 Val Ile Phe Leu Leu Cys Phe Val Pro Tyr Asn Ser Thr Leu Ala Val
201      245      250      255
202 Tyr Gly Leu Leu Arg Ser Lys Leu Val Ala Ala Ser Val Pro Ala Arg
203      260      265      270
204 Asp Arg Val Arg Gly Val Leu Met Val Met Val Leu Leu Ala Gly Ala
205      275      280      285
206 Asn Cys Val Leu Asp Pro Leu Val Tyr Tyr Phe Ser Ala Glu Gly Phe
207      290      295      300
208 Arg Asn Thr Leu Arg Gly Leu Gly Thr Pro His Arg Ala Arg Thr Ser
209      305      310      315      320
210 Ala Thr Asn Gly Thr Arg Ala Ala Leu Ala Gln Ser Glu Arg Ser Ala
211      325      330      335
212 Val Thr Thr Asp Ala Thr Arg Pro Asp Ala Ala Ser Gln Gly Leu Leu
213      340      345      350
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221 <212> TYPE: DNA
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226 gcacacgcgg gactgcgcga cgcgtcttac ctggcgcacc tgtgcgtcgt ggacctcgtg 180
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247 <213> ORGANISM: Homo sapiens
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